Changed the margins in cases where the sequence text was "wrapped" down to the next line.  Edited a format error in the Current Application Data section, specifically:  Edited the Current Application Data section with the actual current number. The number inputted by the applicant was   the prior application data; or   other    Added the mandatory heading and subheadings for "Current Application Data".  Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer. Changed the spelling of a mandatory field (the headings or subheadings), specifically:  Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:  Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: //3  Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.  Inserted colons after headings/subheadings. Headings edited included:  Deleted:   non-ASCil "garbage" at the beginning/end of files;   secretary initials/filename at end of file page numbers throughout text;   other invalid text, such as    Inserted mandatory headings, specifically:  Corrected an obvious error in the response, specifically:  Edited identifiers where upper case is used but lower case is required, or vice versa.  Corrected an error in the Number of Sequences field, specifically:  A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.  Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length: field accordingly (error	umber: _	,	_	ted by th STI		CRF Processin	g Date:_	
Edited the Current Application Data section, specifically:  Edited the Current Application Data section with the actual current number. The number inputted by the applicant was   the prior application data; or   other    Added the mandatory heading and subheadings for "Current Application Data".  Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer. Changed the spelling of a mandatory field (the headings or subheadings), specifically:  Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:  Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: //3  Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.  Inserted colons after headings/subheadings. Headings edited included:  Deleted extra, invalid, headings used by an applicant, specifically:  Deleted:   non-ASCil "garbage" at the beginning/end of files;   secretary initials/filename at end of file page numbers throughout text;   other invalid text, such as    Inserted mandatory headings, specifically:  Edited identifiers where upper case is used but lower case is required, or vice versa.  Corrected an error in the Number of Sequences field, specifically:  A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.  Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error use to a Patentth bug). Sequences corrected:    Other:	Changed	a file from non-A	SCII to ASCII	INTER	ED			(STIC s
Edited the Current Application Data section with the actual current number. The number inputted by the applicant was the prior application data; or other						d" down to the	next line.	
applicant was	Edited a fo	ormat error in the	Current Applic	ation Data section	, specifica	Jly:	· · · · · ·	
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Deleted: non-ASCil "garbage" at the beginning/end of files; secretary initials/filename at end of fi page numbers throughout text; other invalid text, such as Inserted mandatory headings, specifically:	Inserted c	olons after headi	ngs/subheading	gs. Headings edit	ed include	<b>d:</b> , .	•	
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Deleted <b>ending</b> stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a Patentin bug). Sequences corrected:  Other:	A "Hard P	•					deleted.	
Other:								
	Other		•				-	

\*Examiner: The above corrections must be communicated to the applicant in the first Offic Action. DO NOT send a copy of this form.

3/1/95



OIPE

RAW SEQUENCE LISTING DATE: 02/06/2002 PATENT APPLICATION: US/09/905,348 TIME: 08:21:36

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3 <110> APPLICANT: Genentech, Inc.

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              Ferrara, Napoleone
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              Gao, Wei-Qiang
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RAW SEQUENCE LISTING DATE: 02/06/2002 PATENT APPLICATION: US/09/905,348 TIME: 08:21:36

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                    165
                                         170
286 Cys Gln Gln Ala Glu Cys Pro Gly Gly Cys Arg Asn Gly Gly Phe Cys
287
                                     185
                                                         190
                180
289 Asn Glu Arg Arg Ile Cys Glu Cys Pro Asp Gly Phe His Gly Pro His
                                 200
292 Cys Glu Lys Ala Leu Cys Thr Pro Arg Cys Met Asn Gly Gly Leu Cys
                            215
295 Val Thr Pro Gly Phe Cys Ile Cys Pro Pro Gly Phe Tyr Gly Val Asn
296 225
298 Cys Asp Lys Ala Asn Cys Ser Thr Thr Cys Phe Asn Gly Gly Thr Cys
299
                    245
                                         250
301 Phe Tyr Pro Gly Lys Cys Ile Cys Pro Pro Gly Leu Glu Gly Glu Gln
                260
                                     265
304 Cys Glu Ile Ser Lys Cys Pro Gln Pro Cys Arg Asn Gly Gly Lys Cys
305
            275
                                 280
307 Ile Gly Lys Ser Lys Cys Lys Cys Ser Lys Gly Tyr Gln Gly Asp Leu
308
        290
                            295
                                                 300
310 Cys Ser Lys Pro Val Cys Glu Pro Gly Cys Gly Ala His Gly Thr Cys
                        310
                                             315
313 His Glu Pro Asn Lys Cys Gln Cys Gln Glu Gly Trp His Gly Arg His
```

Use of n and/or Xaa has been detected in the Sequence Listing.

Review the Sequence Listing to insure a corresponding explanation is presented in the <220> to <223> fields of each sequence using n or Xaa.

**VERIFICATION SUMMARY** 

DATE: 02/06/2002

PATENT APPLICATION: US/09/905,348

TIME: 08:21:37

Input Set : N:\jumbos\905348.txt

Output Set: N:\CRF3\02062002\1905348.raw

L:34 M:270 C: Current Application Number differs, Replaced Current Application Number L:35 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:511 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 L:512 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 L:513 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 L:514 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 L:769 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26

L:1701 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:50 L:3586 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:113 L:4040 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:131

L:5344 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:174 L:5479 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:175

L:6540 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:206